```
US-09-764-876-20
; Sequence 20, Application US/09764876
; Publication No. US20030044890A1
; GENERAL INFORMATION:
APPLICANT: Rosen et al.
TITLE OF INVENTION: Nucleic Acids, Proteins, and Antibodies
; FILE REFERENCE: PTZ25
; CURRENT APPLICATION NUMBER: US/09/764,876
; CURRENT FILING DATE: 2001-01-17
; PRIOR APPLICATION NUMBER: 60/179,065
; PRIOR FILING DATE: 2000-01-31
; PRIOR APPLICATION NUMBER: 60/180,628
; PRIOR FILING DATE: 2000-02-04
; PRIOR APPLICATION NUMBER: 60/214,886
; PRIOR APPLICATION NUMBER: 60/214,886
; PRIOR APPLICATION NUMBER: 60/217,487
; PRIOR APPLICATION NUMBER: 60/217,487
; PRIOR APPLICATION NUMBER: 60/225,758
; PRIOR FILING DATE: 2000-07-11
; PRIOR APPLICATION NUMBER: 60/225,758
; PRIOR FILING DATE: 2000-08-14
; PRIOR APPLICATION NUMBER: 60/220,963
; PRIOR FILING DATE: 2000-07-26
; PRIOR APPLICATION NUMBER: 60/217,496
; PRIOR APPLICATION NUMBER: 60/217,496
; PRIOR APPLICATION NUMBER: 60/225,447
; PRIOR FILING DATE: 2000-07-11
; PRIOR APPLICATION NUMBER: 60/225,447
; PRIOR FILING DATE: 2000-08-14
```

```
32.9%; Score 2590.5; DB 11; Length 528; 94.3%; Pred. No. 6.1e-221;
Query Match
Best Local Similarity
Matches
   514; Conservative
            2; Mismatches
                    Indels
                       21; Gaps
   Qу
Db
   Qy
Db
   Οy
Db
   Qу
Db
   Oy
Db
     TKPSLTQKEIAKATSIVAWLAKQHNAYGGFSSTQDTVVALQALAKYATTAYMPSEEINLV 1285
Qy
Db
   Qy
Db
   Qу
Db
   Qу
   1466 LSQYT 1470
Qу
Db
   520 TIQYS 524
```

```
US-09-764-853-583
; Sequence 583, Application US/09764853
; Patent No. US20020090672A1
; GENERAL INFORMATION:
; APPLICANT: Rosen et al.
; TITLE OF INVENTION: Nucleic Acids, Proteins, and Antibodies
; FILE REFERENCE: PJZ06
; CURRENT APPLICATION NUMBER: US/09/764,853
; CURRENT FILING DATE: 2001-01-17
; Prior application data removed - consult PALM or file wrapper
; NUMBER OF SEQ ID NOS: 939
; SOFTWARE: Patentin Ver. 2.0
; SEQ ID NO 583
; LENGTH: 531
; TYPE: PRT
; ORGANISM: Homo sapiens
US-09-764-853-583
```

```
33.0%; Score 2596.5; DB 9; Length 531; 94.3%; Pred. No. 1.8e-221; tive 2; Mismatches 8; Indels 21;
Query Match
Best Local Similarity
    516; Conservative
                          21; Gaps
    Qу
Db
     Qy
     SAFGERDGNGNTWLTAFVTKCFGQAQKFIFIDPKNIQDALKWMAGNQLPSGCYANVGNLL 1103
Qy
   1044
Db
   Qу
Db
    Qу
Db
     Qу
   Qy
Db
   Qу
Db
     Qу
Db
   1464 FKLSQYT 1470
Qу
    521 QATIQYS 527
Db
```